Jesse Eaton

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Experience

Senior Machine Learning Scientist, Freenome

September 2019 - Present

- Increased classification performance to industry high of 0.9 sensitivity at 0.9 specificity for detection of colorectal cancer by creating Freenome's core ML model
- Promoted iterations of important models by developing framework for model comparison
- Enabled explosive model development by refactoring and productionizing core custom model
- Streamlined user model reports by redesigned reporting and model diagnostic infrastructure
- Managed machine learning science intern leading to development of robust kmer model pipeline
- Increased performance relative to Protein and DNA signals by identifying RNA markers

Machine Learning Research Engineer, Qeexo

February 2018 - April 2019

- Generated multiple experimental company products by fusing sensor data with efficient machine learning classifiers https://www.youtube.com/watch?v=1S6irWv8G20
- Compressed gradient boosting classifier to 20 kB size for highly time/space/energy constrained environments achieving < 5ms classification time
- Increased customer awareness by demoing projects at Consumer Electronics Show 2019

Graduate ML Researcher, Carnegie Mellon

January 2017 - January 2018

- Developed theory for tumor deconvolution/phylogenetic inference with inclusion of structural variants
- Led design and developed machine learning pipeline for predicting tumor progression
- Accepted as presenter at ISMB 2018 Comp Bio conference for deconvolution paper and published in the journal "Bioinformatics"
 https://academic.oup.com/bioinformatics/article/34/13/i357/5045780

Software Systems Engineer, MITRE

September 2015 - August 2016

 Built web based electronic medical validation tool for Health Services Dept. as main engineer using Amazon Elastic Compute Cloud (AWS)

Leadership and Awards

Culture Leader, Freenome

January 2022

• Nominated as one of 10 company culture leader for exemplifying Freenome's values

Patent for signal discovery, Freenome

April 2020

 Discovered novel RNA signal relating to 11 sets of specific miRNAs by building classifiers for colon cell proliferative disorders

Education

Carnegie Mellon, M.S. Computational Biology

September 2016 - December 2017

• GPA: 3.91, Courses: machine learning, statistics, simulation, regression, computational genomics, automation of biology, cancer biology

Tufts University, B.S. Biomedical Engineering

September 2011 - May 2015

• GPA: 3.45, Courses: genetics, algorithms, quantum chemistry, medical imaging, machine programming and assembly, drug delivery, tissue engineering